

## Sequence Listing

### (1) SEQ ID NO:1

(A) Length: 2096 base pairs

(B) Type: nucleic acid

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1 ctctctgct cctegaagaa ggccagggcg gggctgccgc aagttttgac atttgcag
61 cggagacgcg cgcgggcact ctcgggccga cggtgcggc ggccggcgac cctccagagc
121 ccttagtcg cgccecgcc ctcctctgc cggagtcgc gggccacga ggccecgcc
181 cgtctcccg cgttgcctc cccggcgcc gcagccatgt cccggggcc cgaggaggtg
241 aaccggtca cggagagcac ctaccggaat gttatggaac agttcaatc tgggtgcga
301 aatttaataa acctggggaa aaattatgag aaagtgtaa acgtatgat cctggcagga
361 aaagcctact acgatggagt ggccaagatc ggtgagattg ccaactgggtc cccctgtca
421 actgaactgg gacatgtcct catagagatt tcaagtacc acaagaaact caacgagagt
481 ctgatgaaa attttaaaaa attcacaaa gagattatcc atgagctgga gaagaagata
541 gaacttgacg tgaaatatat gaacgcaact ctaaaagat accaaacaga acacaagaat
601 aaattagagt cttggagaa atccaagct gagtgaaga agatcagaag gaaaagccaa
661 ggaagccgaa acgcactcaa atatgaacac aaagaaattg agtatgtgga gaccgttact
721 tctctcaga gtgaaatcca gaaattcatt gcagatggtt gcaaagagc tctgctttaa
781 gagaagagc gcttctgctt tctggtgat aagcactgtg gctttgcaa ccacatacat
841 tattactact tacagtctgc agaactactg aattccaagc tgcctcggtg gcaggagacc
901 tgtgttgatg ccatcaaagt gccagagaaa atcatgaata tgatcgaaga aataaagacc
961 ccagcctcta cccctgtgc tggaaactct caggttcac ccatgatcga gagaagcaat
1021 gtggttagga aagattacga cacccttct aatgtctac caaatgccc cccctctct
1081 tcaggcagag catataccag tccttgatc gatatttta ataaccagc caggtgcgc
1141 ccgaattcac aaagggtaaa taattcaaca ggtacttcg aagatcccag ttacagcga
1201 tcagtttcgg ttgcaacggg actgaacatg atgaagaagc agaaagtga gaccatctc
1261 cgcacactg cgggtccaa caagacctt ctcagcttg cacagggaga tgcacacg
1321 ctgctatcc cggaggagaa ggtggttg ctctatggag aacacgacgt gtccaaggcg
1381 aggggttggt tccgtctgc gtacacgaag ttgtggaag aaaatgagac agaagcagt
1441 accgtgccc cgcgaagccc cacaccagtg agaagcatc gcaccgtgaa cttgtctgag
1501 aatagcagtg ttgtatccc cccaccgac tacttggaat gctgtccat gggggcagct
1561 gccagacgga gagcagatc ggccaggacg acatccact ttaaggcccc agcgtccaag
1621 cccgagaccg cggctctaa cgtgccaac gggactgcaa agcgccttt tctcagcga
1681 gaaaaccct ttgccactg gaaactcgc cgcactgtg cgaatgatc ctcggcacc
1741 atcattcgat gagaggacag ccaaggactc tccgggct ctcgggttct ccttgcgga
1801 atgatggcg cactctgtc gccactgct gacggtggg aagcttcagt ggagaggcct
1861 aactctaag tgcctgctt aagcaaatc tcttctctg ttacagtag ttgggtgac
1921 aagtttctc cttaagata aatgagtaat agtctaata ccagctcagc catttaaat
1981 attttcttc tattctgtc aagaaacagt aaacttggt tcaatctta aaaaaaaaaa
2041 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaa

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### (2) SEQ ID NO:2

(A) Length: 511 amino acids

## (B) Type: amino acid

1 msrgpeevnr ltestyrnvm eqfnpglrnl inlgknyeka vnamilagka yydgvakige  
 61 iatgspvste lghvlieiss thkklnesld enfkkfhkei ihelekkie dvkymnatlk  
 121 ryqtehnkl esleksqael kkirrksgs rnalkyehke ieyvetvtsr qseiqkfiad  
 181 gckealleek rrfcflvdkh cgfanhihy hlqsaellns klprwqetcv daikvpekim  
 241 nmieektpa stpvsqtpqa spmiersnvv rkdydtlskc spkmppapsg raytsplidm  
 301 fnnpataapn sqrvnnstgt sedpslqrsv svatglnmmk kqkvktifph tagsnktlls  
 361 faqgdvitll ipeekdgwly gehdvskarg wfpssytkll eeneteavtv ptpsptpvr  
 421 istvnlsens svvipppdyl eclsmgaaad rradsartts tfkapaskpe taapndangt  
 481 akppflngen pfatvklrpt vtndrsapii r

## (3) SEQ ID NO:3

(A) Length: 1794 base pairs

(B) Type: nucleic acid

1 cacatataat aatagcaact cctggtegac tgattgacca ctgaaacct ttegtatttt  
 61 ccaagtgetg gcaagegett cctgegcagg cegaggegac ctggagtttg tgacgetgtg  
 121 atggtctaga ggettgagat tcaagatctg ggtgceatca tttctggtt ctgttgatga  
 181 cctcttcca ggttacatag agettacatc ttgcatcctc aagcgttttt cttataagge  
 241 taaaaattca caaagcatat atcaatgaat caggaggatc tagatccgga tagtactaca  
 301 gatgtgggag atgttacaaa tactgaagaa gaacttatta gagaatgtga agaaatgtgg  
 361 aaagatatgg aagaatgtca gaataaatta tcactattg gaactgaaac actcaccgat  
 421 tcaaatgtc agctatcatt gtaattatg caagtaaaat gttaaccgc tgaactcagt  
 481 caatggcaga aaaaaacacc tgaacaatt ccttgactg aagacgttct cataacatta  
 541 ggaaaagaag agttcaaaaa gctgagacaa gatcttgaaa tggtagtgc cactaaggag  
 601 tcaagaatg aaaagttaa ggaagactta gaaagggaa aacgggtggt ggatgaacag  
 661 caacagataa tggaatctct taatgtacta cacagtgaat tgaataataa gggtgaaaca  
 721 tttctgaat caagaattct taatgaactg aaaactaaaa tgettaatat aaaagaatat  
 781 aaggagaaac tcttgagtac ctggggcgag ttctagaag accattttcc tctgcctgat  
 841 agaagtgtta aaaagaaaaa gaaaaacatt caagaatcat ctgtaacct gataacactg  
 901 catgaaatgt tagagattct tataataga ttatttgatg ttccacatga tccatatgct  
 961 aaaattagtg attccttttg gccacettat gttgagctgc tctgcgtaa tgggaattgcc  
 1021 ttgagacatc cagaagatcc aaccgaata agattagaag ctttccatca gtaaaaggat  
 1081 gttttctttt ttacacagt aaaaattctt atcattcaag gatattggaa ccacaggact  
 1141 atttgataa aaaacattat ttgcaatta atgcgcatag tacttttatt gcaaaatggc  
 1201 atgtgetgcc atctattatt catttttaa tggcatttc ttattcagt agtgettttag  
 1261 tgttttaaac tatatggata agaattgcagg tagataatat tctaggcata aaacatttaa  
 1321 tgtaccttac ctcattgaat attctttgga ttctttgttg atttatgata ttgctaatat  
 1381 aatattttct taaaatatat aacaatatct ttatgcatt tgagtccag ctgggtgcttc  
 1441 tttatattta gaaattataa tgggaaggct atttaatta cagatgggtt taaaattgag  
 1501 gtaatatctg aggtggcata atttaaaaat atttagcaaa ttgtttcat atatactgtc  
 1561 ttatttctag attgttttaa aattggaata tgaataacta atggataaag ctgcataaa  
 1621 attgatattt tagttgttat tattaatata tcatgttacc ttatatatta atctactctt  
 1681 gattctgcta attattacca acaaaattgt attcatgaca ttttattaat cctctgtgaa  
 1741 tttctgtaa ataaaattat ttctgaaat ctctaaaaaa aaaaaaaaaa aaaa

(4) SEQ ID NO:4

(A) Length: 269 amino acids

(B) Type: amino acid

1 mnqedldpds ttdvgdvtnt eelirecee mwkdmeecqn klsligteti tdsnaqlsl  
61 imqvkcltae lsqwqkktp tipltedvli tlgkeefqkl rqllemvlt keskneklke  
121 dlereqrwld eqqqimesln vlhseknkv etfsesrfin elktkmlnik eykekllstl  
181 gefledhfpl pdrsvkkkkk niqessvnl tthemleili nrlfdvphdp yvkisdsfwp  
241 pyvelllrng ialrhpedpt rirleafhq

(5) SEQ ID NO:5

(A) Length: 1660 base pairs

(B) Type: nucleic acid

1 gccaggaaat gccctggagt gtgtgtcacc tgtccaggac gacttgttga tteccaggag  
61 ggccgcttt cgggtctggg tccccgagag gactgccttg ctcacctgac cctcggegc  
121 ggccccgggg agctcccag aggcceccgg gatcgctggc cctccgaact ccacagcaat  
181 gagcaagttg ggcaagttct ttaaaggggg cggtcttct aagagccgag ccgtccag  
241 tccccaggag gccctggtec gacttcggga gactgaggag atgctgggca agaaacaaga  
301 gtacctggaa aatgaatcc agagagaaat cgcctggcc aagaagcac gcacgcagaa  
361 taagcgagct gcattacagg cactaaagag aaagaagagg ttcgagaaac agctcacta  
421 gattgatggc acatttcta ccattgagtt ccagagagaa gccctggaga actcacacac  
481 caacactgag gtgttgagga acatgggctt tgcagcaaaa gcgatgaaat ctgttcata  
541 aaacatggat ctgaacaaaa tagatgatt gatgcaagag atcacagagc aacaggatat  
601 cgccecaaga atctcagaag cattttctca acgggttggc tttggtgatg actttgatga  
661 ggatgagttg atggcagaac ttgaagaatt ggaacaggag gaattaaata agaagatgac  
721 aaatatecgc ctcccaaatg tgccttctc ttctctcca gcacagccaa atagaaaacc  
781 aggcagtgcg tccactgcac gtcgacccg agcagcatct tcccagaggc cagaagaaga  
841 ggatgatgat atcaacaat tggcagcttg ggctacctaa actaaaacac attttgata  
901 cctaaattaa tgagctatag ataaaatata aaaaatgttt ttaccaagtt cagaagttaa  
961 caaagactct gctttataat tatattgaat gaataattgt gtttaagcc tctaagtaa  
1021 aagtaaaaaa ggagtcattg gcatacatag aatcagtgat ggaggccagg caccgtatct  
1081 catgcctata atcccagcac tttgggaggg tgaggcagtt gagaccagga gttcgagtc  
1141 agcctgacca acatgaagaa accctgtctg tactaaaaat acaaaaatta gccggacatg  
1201 gtggcaggca cctgtaatcc cagctacttg ggaggctgag tcaggagaat cgcttgagcc  
1261 caggagatgg aggttgcatg gagccaagat catgccactg cactccagac tgggcaacag  
1321 agggagactc cgtctcaaaa actaaaaaaa aaaaatacat ttagtatagc ggggggtggg  
1381 ggggagaaat aatgttattt cctatgcgaa tgacgtgtat cctgtacc atggtaaatg  
1441 taaatatact gtgtctcttt tgggagagcc ttttagtaga ggagtcttat atgagtctt  
1501 acataagtag ttctacttga gttttcagtt ttgaaatctt aaaggagctt taattgacat  
1561 ttattatacc aattaagctt ggaatggggc aatggatgca ttcccaaaa cgtgtgaaag  
1621 cactaacagc ttattattgt gaatgagaat ctctgggtg

(6) SEQ ID NO:6

(A) Length: 233 amino acids

(B) Type: amino acid

1 msklgkffkg ggssksraap spqealvrlr eteemlgkkq eylenriqre ialakkhgtq  
 61 nkraalqalk rkkrfekqlt qidgtlstie fqrealensh ntevlrnmg faakamksvh  
 121 enmdlknidd lmqeiteqqd iaqaiseafs qrvfgddfd edelmaelee leqeelnkkm  
 181 tnirlpnvps sslpaqpnrk pgmsstarrs raassqraee edddikqlaa wat

(7) SEQ ID NO:7

(C) Length: 1666 base pairs

(D) Type: nucleic acid

1 agcgggcccgcg gggctgcgag ctagggcggg agaaggagcg cggggaggac gtacctgtg  
 61 agatgcgagc cggccaacag cttgcaagca tgcctcgtg gaccgcagcc tggaggctcc  
 121 cgcgtgaggg actegggccc cagggcccta gcttcgcgag ggtgcctgtc gcaccagca  
 181 gcagcagcgg cggccgaggg ggcgcgcgagc cgaggccgct tccgtttcc tacaggcttc  
 241 tggacgggga ggcagccctc cggccgctgc tctttttgca cgggtcttc ggcagcaaaa  
 301 ctaactcaa ctccatgcc aagatcttgg ccagcagac aggcctgct gacggtggat  
 361 gctcgtaac acggtgacag cccccacag ccagacatga gctacgagat catgagccag  
 421 gacctgcagg accttctgcc ccagctgggc ctggtgcct gcgtcgtct tggccacagc  
 481 atgggaggaa agacagccat gctgctggca ctacagaggg tgagccgcc atgtctgggg  
 541 cctctccca ttcagtatat acctgaggg cctgcagge aacctgggac tcacatgac  
 601 gttgatgac caagttcagg ctccaggagc catgctgag actccctatg tctgctaag  
 661 actggtccca gttcggttct ctccacagc cagagctgg ggaacgtctc attgctgtag  
 721 atatcagccc agtggaagc acaggtgtct cccacttgc aacctatgtg gcagccatga  
 781 gggccatcaa catcgcagat gagctgccc gctccgtgc cggaaaactg gcggatgaac  
 841 agtcagttc tgtcatccag gacatggccg tgcggcagca cctgtcact aacctggtag  
 901 aggtagacgg gcgttctgtg tggagggtga acttggtatg cctgaccag cacctagaca  
 961 agatcttggc ttccacag aggcaggagt cctacctcg gccaacactc ttctctctg  
 1021 gtggaaactc ccagttctg catccagcc accacctga gattatcgg ctcttcctc  
 1081 gggcccagat gcagacggtg cgaacgtg gccactggat ccacgtgac cggccacagg  
 1141 actcatage tgccatcga ggttctctg tctaagagt gctggcaaga agatggcgg  
 1201 gcgtgtgtgc tcatgctgt aattccagca ctttgggagg ctaaggcggg aggatgactt  
 1261 gagccagga gttggagacc agcctggcca acatggtgaa accctgtct tactaaaaat  
 1321 acaaaaatta gcttggtg gtgtgcaca cctgtaatc cagctactct ggaggctgag  
 1381 gcaggagaat cacttgaacc ctggaggcag aggttgcaat gagccgagat cacaccacta  
 1441 cactccagcc taggcaacag agcaagactc tgtctcaaaa aaacaaaac aaaaaggagg  
 1501 cacaiaaacc caggttcaa gtctctgcag cctgtccac atttgggcac agaaggactc  
 1561 agacaggcac tgtgtgggca cgaggttcta cagggtgtg cagacctcag gctttaatga  
 1621 ataaagacac tactcccaa aaaaaaaaaa aaaaaaaaaa aaaaaa

(8) SEQ ID NO:8

(A) Length: 101 amino acids

(B) Type: amino acid

1 mragqqlasm lrwtrawrlp reglgphgps farvpvapss ssggrggaep rplplsyrll  
 61 dgeaalpavv flhglfgskt nfnsiakila qqtgradggc s

(9) SEQ ID NO:9

(A) Length: 23 base pairs

(B) Type: nucleic acid

GCTTCTGCTTTCTGGTTGATAAG

(10) SEQ ID NO:10

(A) Length: 22 base pairs

(B) Type: nucleic acid

GATTTTCTCTGGCACTTTGATG

(11) SEQ ID NO:11

(A) Length: 22 base pairs

(B) Type: nucleic acid

GAAACACTCACCGATTCAAATG

(12) SEQ ID NO:12

(A) Length: 22 base pairs

(B) Type: nucleic acid

GCTTTTGGAACCTTCTTTTCC

(13) SEQ ID NO:13

(A) Length: 22 base pairs

(B) Type: nucleic acid

AGGGCAGAAGAAGAGGATGATG

(14) SEQ ID NO:14

(A) Length: 22 base pairs

(B) Type: nucleic acid

GTGCTGGGACTATAGGCATGAG

(15) SEQ ID NO:15

(A) Length: 20 base pairs

(B) Type: nucleic acid

CTGACGGTGGATGCTCGTAA

(16) SEQ ID NO:16

(A) Length: 22 base pairs

(B) Type: nucleic acid

GTGCCTGTCTGAGTCCTTCTGT

(17) SEQ ID NO:17

(A) Length: 21 base pairs

(B) Type: nucleic acid

AAGCACTGTGGCTTTGCAAAC

(18) SEQ ID NO:18

(A) Length: 21 base pairs

(B) Type: nucleic acid

AACACTCACCGATTCAAATGC